

SEOUENCE LISTING

<110> Daniel E.H. Affilia
Arthur B. Raitano
Rene S. Hubert
Steve Chappell Mitchell
Aya Jakobovits

<120> NOVEL GENE UPREGULATED IN CANCERS OF THE PROSTATE

<130> 129.21-US-U1

<140> 09/697,206

<141> 2000-10-26

<150> 60/162,364 <151> 1999-10-28

<160> 26

<170> FastSEQ for Windows Version 4.0

<210> 1

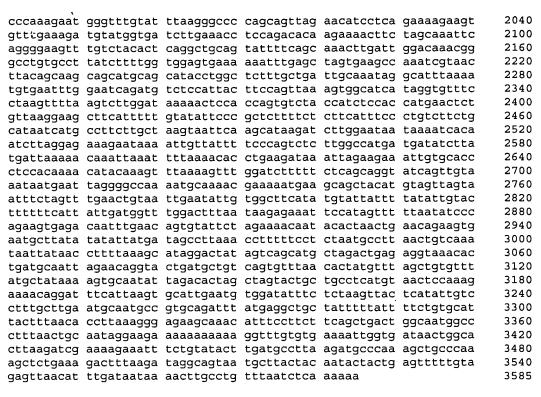
<211> 3585

<212> DNA

<213> Homo Sapiens

<400> 1

ctttttggga tcactgctgg ggccaccggg gccaagctag gctcggatga gaaggagttg 60 atcctgctgt tctggaaagt cgtggatctg gccaacaaga aggtgggaca gttgcacgaa 120 gtgctagtta gaccggatca gttggaactg acggaggact gcaaagaaga aactaaaata 180 gacgtcgaaa gcctgtcctc ggcgtcgcag ctggaccaag ccctccgaca gtttaaccag 240 300 tcagtgagca atgaactgaa tattggagta gggacttcct tctgtctctg tactgatggg cagetteatg teaggeaaat cetgeateet gaggetteea agaagaatgt actattacet 360 gaatgettet atteettttt tgatettega aaagaattea agaaatgttg eeetggttea 420 cctgatattg acaaactgga cgttgccaca atgacagagt atttaaattt tgagaagagt 480 agttcagtct ctcgatatgg agcctctcaa gttgaagata tggggaatat aattttagca 540 atgatttcag agccttataa tcacaggttt tcagatccag agagagtgaa ttacaagttt 600 660 gaaagtggaa cttgcagcaa gatggaactt attgatgata acaccgtagt cagggcacga ggtttaccat ggcagtcttc agatcaagat attgcaagat tcttcaaagg actcaatatt 720 780 gccaagggag gtgcagcact ttgtctgaat gctcagggtc gaaggaacgg agaagctctg gttaggtttg taagtgagga gcaccgagac ctagcactac agaggcacaa acatcacatg 840 gggacccggt atattgaggt ttacaaagca acaggtgaag atttccttaa aattgctggt 900 ggtacttcca atgaggtagc ccagtttctc tccaaggaaa atcaagtcat tgttcgcatg 960 cqqqqqctcc ctttcacqqc cacaqctqaa qaaqtqqtqq ccttctttqq acaqcattqc 1020 cctattactg ggggaaagga aggcatcctc tttgtcacct acccagatgg taggccaaca 1080 1140 ggggacgctt ttgtcctctt tgcctgtgag gaatatgcac agaatgcgtt gaggaagcat aaagacttgt tgggtaaaag atacattgaa ctcttcagga gcacagcagc tgaagttcag 1200 caggigetga ategatiete eteggeeeet eteaticeae tiecaaeeee teecatiati 1260 ccagtactac ctcagcaatt tgtgccccct acaaatgtta gagactgtat acgccttcga 1320 ggtcttccct atgcagccac aattgaggac atcctggatt tcctggggga gttcgccaca 1380 qatattcqta ctcatqqqqt tcacatqqtt ttqaatcacc aqqqccqccc atcaqqaqat 1440 gcctttatcc agatgaagtc tgcggacaga gcatttatgg ctgcacagaa gtgtcataaa 1500 aaaaacatga aggacagata tgttgaagtc tttcagtgtt cagctgagga gatgaacttt 1560 gtgttaatgg ggggcacttt aaatcgaaat ggcttatccc caccgccatg cctgtctcct 1620 ccctcctaca catttccagc tcctgctgca gttattccta cagaagctgc catttaccag 1680 ccctctgtga ttttgaatcc acgagcactg cagccctcca cagcgtacta cccagcaggc 1740 acteagetet teatgaacta cacagegtae tateecagee eeccaggtte geetaatagt 1800 cttggctact teectacage tgctaatett ageggtgtee etccacagee tggcacggtg 1860 gtcagaatgc agggcctggc ctacaatact ggagttaagg aaattcttaa cttcttccaa 1920 ggttaccagt atgcaaccga ggatggactt atacacacaa atgaccaggc caggactcta 1980



<210> 2 <211> 517 <212> PRT <213> Homo Sapiens

<400> 2

10 Gly Ala Ser Gln Val Glu Asp Met Gly Asn Ile Ile Leu Ala Met Ile 25 Ser Glu Pro Tyr Asn His Arg Phe Ser Asp Pro Glu Arg Val Asn Tyr Lys Phe Glu Ser Gly Thr Cys Ser Lys Met Glu Leu Ile Asp Asp Asn Thr Val Val Arg Ala Arg Gly Leu Pro Trp Gln Ser Ser Asp Gln Asp 75 Ile Ala Arg Phe Phe Lys Gly Leu Asn Ile Ala Lys Gly Gly Ala Ala 90 Leu Cys Leu Asn Ala Gln Gly Arg Arg Asn Gly Glu Ala Leu Val Arg 105 Phe Val Ser Glu Glu His Arg Asp Leu Ala Leu Gln Arg His Lys His 115 120 125 His Met Gly Thr Arg Tyr Ile Glu Val Tyr Lys Ala Thr Gly Glu Asp 130 135 140 Phe Leu Lys Ile Ala Gly Gly Thr Ser Asn Glu Val Ala Gln Phe Leu 150 155 Ser Lys Glu Asn Gln Val Ile Val Arg Met Arg Gly Leu Pro Phe Thr 170 Ala Thr Ala Glu Glu Val Val Ala Phe Phe Gly Gln His Cys Pro Ile 185 Thr Gly Gly Lys Glu Gly Ile Leu Phe Val Thr Tyr Pro Asp Gly Arg 200 Pro Thr Gly Asp Ala Phe Val Leu Phe Ala Cys Glu Glu Tyr Ala Gln Asn Ala Leu Arg Lys His Lys Asp Leu Leu Gly Lys Arg Tyr Ile Glu

Met Thr Glu Tyr Leu Asn Phe Glu Lys Ser Ser Ser Val Ser Arg Tyr

<212> DNA





```
235
225
                  230
Leu. Phe Arg Ser Thr Ala Ala Glu Val Gln Gln Val Leu Asn Arg Phe
            245
                       250
Ser Ser Ala Pro Leu Ile Pro Leu Pro Thr Pro Pro Ile Ile Pro Val
          260
                               265
Leu Pro Gln Gln Phe Val Pro Pro Thr Asn Val Arg Asp Cys Ile Arg
                           280
Leu Arg Gly Leu Pro Tyr Ala Ala Thr Ile Glu Asp Ile Leu Asp Phe
                       295
                                          300
Leu Gly Glu Phe Ala Thr Asp Ile Arg Thr His Gly Val His Met Val
                  310
                                   315
Leu Asn His Gln Gly Arg Pro Ser Gly Asp Ala Phe Ile Gln Met Lys
                                 330
              325
Ser Ala Asp Arg Ala Phe Met Ala Ala Gln Lys Cys His Lys Lys Asn
                              345
Met Lys Asp Arg Tyr Val Glu Val Phe Gln Cys Ser Ala Glu Glu Met
                  360
Asn Phe Val Leu Met Gly Gly Thr Leu Asn Arg Asn Gly Leu Ser Pro
                       375
Pro Pro Cys Leu Ser Pro Pro Ser Tyr Thr Phe Pro Ala Pro Ala Ala
                   390
                                      395
Val Ile Pro Thr Glu Ala Ala Ile Tyr Gln Pro Ser Val Ile Leu Asn
Pro Arg Ala Leu Gln Pro Ser Thr Ala Tyr Tyr Pro Ala Gly Thr Gln
                               425
                                                 430
Leu Phe Met Asn Tyr Thr Ala Tyr Tyr Pro Ser Pro Pro Gly Ser Pro
       435
                          440
Asn Ser Leu Gly Tyr Phe Pro Thr Ala Ala Asn Leu Ser Gly Val Pro
                       455
                                          460
Pro Gln Pro Gly Thr Val Val Arg Met Gln Gly Leu Ala Tyr Asn Thr
                   470
                                      475
Gly Val Lys Glu Ile Leu Asn Phe Phe Gln Gly Tyr Gln Tyr Ala Thr
                           490
              485
Glu Asp Gly Leu Ile His Thr Asn Asp Gln Ala Arg Thr Leu Pro Lys
          500
                             505
Glu Trp Val Cys Ile
       515
<210> 3
<211> 14
<212> DNA
<213> Artificial Sequence
<220>
<223> Primer
<400> 3
                                                                     14
ttttgatcaa gctt
<210> 4
<211> 42
<212> DNA
<213> Artificial Sequence
<220>
<223> Adaptor
ctaatacgac tcactatagg gctcgagcgg ccgcccgggc ag
                                                                     42
<210> 5
<211> 12
```

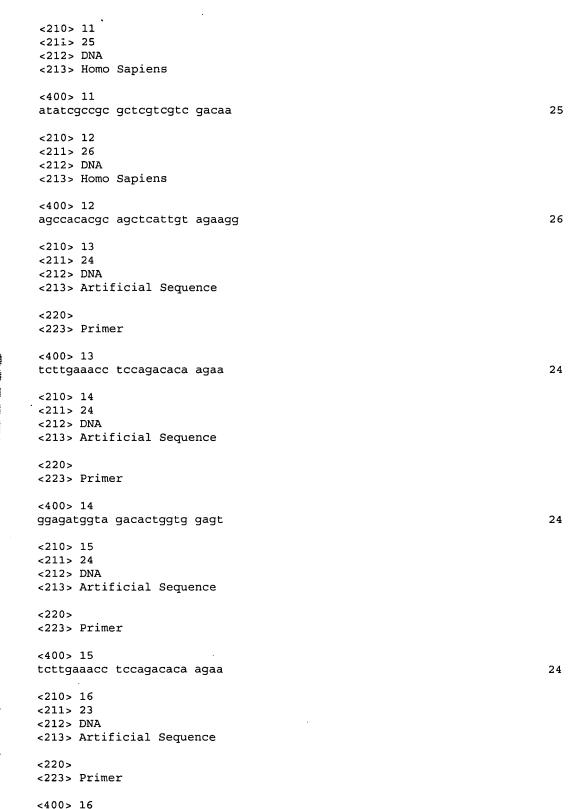




	<213> Artificial Sequence	
	; <220>	
	<223> Adaptor	
	<400> 5	
	ggcccgtcct ag	12
	<210> 6	
	<211> 40	
	<212> DNA <213> Artificial Sequence	
	<220> <223> Adaptor	
	C2237 Adaptor	
	<400> 6	
	gtaatacgac tcactatagg gcagcgtggt cgcggccgag	40
	<210> 7	
	<211> 10 <212> DNA	
	<213> Artificial Sequence	
	<220> <223> Adaptor	
	<400> 7	10
	cggctcctag	10
	<210> 8	
	<211> 22 <212> DNA	
	<213> Artificial Sequence	
	<220> <223> Primer	
	<400> 8	22
	ctaatacgac tcactatagg gc	22
	<210> 9	
	<211> 22 <212> DNA	
	<213> Artificial Sequence	
	<220> <223> Primer	
	<400> 9	22
	tcgagcggcc gcccgggcag ga	22
	<210> 10	
	<211> 20 <212> DNA	
	<213> Artificial Sequence	
	<220> <223> Primer	
'		
	<400> 10 agcqtqqtcq cqqccqaqqa	20
	ugogoggoog oggoogagga	20

aagttacgat ttggcttcac tgg

<210> 17 <211> 9 <212> PRT



23





```
<213> Homo Sapiens
<400> 17
Phe Leu Gly Glu Phe Ala Thr Asp Ile
<210> 18
<211> 9
<212> PRT
<213> Homo Sapiens
<400> 18
Val Leu Phe Ala Cys Glu Glu Tyr Ala
<210> 19
<211> 9
<212> PRT
<213> Homo Sapiens
<400> 19
Leu Leu Gly Lys Arg Tyr Ile Glu Leu
                 5
<210> 20
<211> 9
<212> PRT
<213> Homo Sapiens
<400> 20
Gln Gln Phe Val Pro Pro Thr Asn Val
1
                5
<210> 21
<211> 9
<212> PRT
<213> Homo Sapiens
<400> 21
Cys Ser Ala Glu Glu Met Asn Phe Val
1
<210> 22
<211> 9
<212> PRT
<213> Homo Sapiens
<400> 22
Phe Leu Ser Lys Glu Asn Gln Val Ile
<210> 23
<211> 9
<212> PRT
<213> Homo Sapiens
Ser Leu Gly Tyr Phe Pro Thr Ala Ala
<210> 24
<211> 9
<212> PRT
```